\_\_\_\_\_

\_\_\_\_\_\_

Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2007; month=12; day=21; hr=16; min=56; sec=13; ms=427;]

## Validated By CRFValidator v 1.0.3

Application No: 10622373 Version No: 2.0

Input Set:

Output Set:

**Started:** 2007-12-03 19:02:21.803

**Finished:** 2007-12-03 19:02:24.017

**Elapsed:** 0 hr(s) 0 min(s) 2 sec(s) 214 ms

Total Warnings: 5

Total Errors: 0

No. of SeqIDs Defined: 9

Actual SeqID Count: 9

Error code		Error Description												
W	402	Undefined organism found in <213> in SEQ ID (3)												
W	213	Artificial or Unknown found in <213> in SEQ ID (4)												
W	213	Artificial or Unknown found in <213> in SEQ ID (7)												
W	213	Artificial or Unknown found in <213> in SEQ ID (8)												
W	213	Artificial or Unknown found in <213> in SEQ ID (9)												

## SEQUENCE LISTING

```
<110> The Regents of the University of California
     Whistler, Jennifer L
<120> METHODS AND COMPOSITIONS FOR MODULATING AGONIST-INDUCED
     DOWNREGULATION OF G PROTEIN-COUPLED RECEPTORS
<130> 316E-001510US (12101-011-999)
<140> 10622373
<141> 2003-07-18
<160> 9
<170> PatentIn version 3.1
<210> 1
<211> 4185
<212> DNA
<213> Homo sapiens
<400> 1
     atgactgggg cagagattga gtctggtgcc caggtcaagc ctgaaaagaa gcctggggaa
     gaggttgtag gtggggctga gatagagaat gatgtccctc tggtggtcag acccaaggtt
     aggacccagg cccagataat gcctggggca aggcccaaga ataagtccaa ggttatgcct
     ggagcaagca ccaaagttga gacaagtgca gtgggtgggg cacgccctaa gagtaaggcc
     aaggcaatac ctgtttcacg atttaaggaa gaagcccaga tgtgggctca gcccaggttt
     ggtgctgaaa gattgtctaa gacagagaga aactcccaga ccaatatcat agcctctcca
     cttgtcagta ctgattctgt cttggttgct aaaacaaagt acctgtctga ggatagagaa
     ctggttaata cagacactga gagctttcct agaaggaagg cccattacca agcaggattc
     cagcettett ttaggteaaa ggaggagaee aatatggggt eetggtgetg teetaggeet
     acatccaaac aagaagcctc tcctaattct gatttcaaat gggtagacaa atctgtgagt
     tccttgttct ggagtggaga tgaggtcact gcaaaatttc atcctgggaa tagggtaaaa
     gacagtaaca gatccatgca catggccaat caagaggcta ataccatgtc taggtcccaa
     actaaccagg agctctatat tgcatctagt tctggttctg aggatgagtc tgttaagaca
     ccctggttct gggccagaga taaaaccaat acctggtctg ggcccaggga agatcccaat
     agcaggtcca ggtttaggtc taagaaagaa gtctatgttg aatcaagttc tggatctgag
     catgaagacc atttggagtc ctggtttggg gctggaaagg agggcaaatt caggtccaaa
     atgagagetg ggaaggagge caataacagg gecaggeaca gggeeaageg agaagettge
     attgatttca tgcctgggtc tatagatgta attaaaaaag agtcctgttt ctggcctgaa
     gaaaatgcta ataccttttc aaggcccatg atcaagaaag aggccagggc cagagcaatg
     acaaaggaag aggccaaaac caaggcccga gccagggcca agcaagaagc caggtcagag
     gaggaagccc tcattgggac ctggttctgg gctacagacg agtccagcat ggcagatgaa
     gccagcatag agtccagtct acaagtggag gatgagtcca taattgggag ttggttctgg
     actgaagaag aggccagtat ggggactggg gctagcagta aatccagacc aaggactgat
     ggggagcgta ttggtgattc cttatttggg gctagggaaa agaccagtat gaaaactggg
     gctgaggcca cctctgaatc tatactagca gctgatgatg aacaggtcat tattggttcc
     tggttctggg ctggtgaaga ggtcaaccaa gaggctgagg aagagaccat ttttgggtcg
     tggttctggg tcattgatgc ggccagtgtg gaatctggtg ttggggtcag ctgtgagtcc
     aggacaaggt ctgaggaaga agaggtcatt ggtccctggt tttggtctgg agaacaagtt
     gatatagagg ctggaatcgg agaagaggcc aggccaggag ctgaagaaga gacaatattc
```

gggtcctggt tttgggctga aaaccagacc tatatggatt gtagggctga aactagctgt gacaccatgc aaggggctga ggaggaggag cccattattg ggtcctggtt ttggaccaga

gtagaagctt gtgtggaggg tgatgtcaac agcaagtcta gcctggagga caaggaagag

gccatgatac catgttttgg agccaaagaa gaggtcagta tgaagcatgg gactggtgtc

60

120

180 240

300

360

420

480 540

600

660 720

780 840

900

1020

1080 1140

1200

1260

1320 1380

1440 1500

1560

1620 1680

> 1740 1800

1860

1920 1980

960

```
agatgcagat ttatggcagg ggctgaggag accaataata agtcttgctt ctgggcagaa
                                                                   2040
aaagaaccct gtatgtatcc tgccggtgga ggaagttgga agtctaggcc agaggaggaa
                                                                   2100
gaggacattg tcaattcgtg gttctggtcc agaaaataca caaagccaga ggccattata 2160
gggtcctggt tatgggctac agaagagagt aatatagatg ggactggaga aaaggccaag 2220
ttactgactg aagaggagac cataatcaat tcctggttct ggaaagaaga tgaagccatt
                                                                   2280
tcagaggcta ctgacagaga agagtccagg ccagaagctg aggaggggga cattgttggt 2340
tcttggttct gggctggaga agaggacaga ctagagccag ctgctgagac tagagaagaa
                                                                   2400
gacaggetag cagetgagaa agaaggtatt gttgggteet ggtttgggge cagagaagag
                                                                   2460
accattagaa gagaggctgg gtcttgcagc aaatccagtc ctaaagctga agaggaagaa
                                                                   2520
                                                                   2580
gtcattattg ggtcctggtt ctgggaagaa gaggccagtc cggaggcagt ggcaggagtc
ggctttgagt caaagcctgg gactgaggag gaagaaatca ctgttgggtc ctggttctgg
                                                                   2640
                                                                   2700
cctgaagaag aagccagtat acaggctgga tctcaggcag tagaggaaat ggagtcagag
actgaagagg aaaccatttt tgggtcctgg ttctgggatg gaaaagaagt cagtgaagaa
                                                                   2760
gcaggaccat gctgtgtatc caagccagag gatgatgaag agatgattgt tgagtcctgg
                                                                   2820
                                                                   2880
ttctqqtcta qaqacaaaqc cattaaqqaa actqqaactq tqqccacctq tqaqtccaaq
ccagaaaatg aggaaggggc cattgttggg tcttggtttg aggctgaaga tgaggtagat
                                                                   2940
                                                                   3000
aacaggactg acaatggaag caactgtggg tccaggacat tagctgatga agatgaggcc
                                                                   3060
atagtggggt cctggttctg ggcaggagat gaggcccatt ttgaatcaaa tcctagcccc
gtgttcaggg ccatttgcag gtccacgtgt tcagttgaac aggagcctga tccttcacgc
                                                                   3120
aggcctcaga gttgggagga ggtcactgtt cagttcaagc ctggtccatg gggtagggtc
                                                                   3180
ggcttcccat ctataagccc ctttagattt ccgaaagagg cagcatcttt attctgtgaa
                                                                   3240
atgtttgggg gcaaacccag gaacatggta cttagcccag aaggggaaga tcaggaatct
                                                                   3300
                                                                   3360
ttgcttcagc ctgatcagcc tagtcctgag ttcccatttc agtatgatcc ttcctacagg
tcagtccagg aaattcgaga gcatcttagg gccaaggaga gtacagagcc tgagagttca
                                                                   3420
tcctgtaact gcatacaatg tgagctgaaa attggttctg aagagtttga agaactcctt
                                                                   3480
ttattaatgg aaaaaattcg ggatcctttt attcatgaaa tatctaaaat cgcaatgggt
                                                                   3540
atgagaagtg cttctcaatt tacccgagat ttcattcgag attcaggtgt tgtctcactt
                                                                   3600
attgaaacct tgcttaatta tccgtcctcc cgagttagaa caagtttttt ggaaaatatg
                                                                   3660
attcgcatgg ccccacctta tccgaatcta aacataattc agacatacat atgtaaagtg
                                                                   3720
tgtgaggaaa cccttgctta tagcgtggat tccccggaac agctgtctgg aataaggatg
                                                                   3780
attagacatc tcactactac tactgactat cacacactgg ttgccaatta tatgtctggg
                                                                   3840
tttctctcct tattagctac aggcaatgcc aaaacaaggt ttcatgtttt gaaaatgcta
                                                                   3900
                                                                   3960
ctgaatttgt ctgaaaatct tttcatgaca aaagaactac tcagtgctga agcagtgtca
gaatttatag gcctctttaa cagggaagag acaaatgaca atattcaaat tgttcttgca
                                                                   4020
atatttgaga atattggcaa caatatcaaa aaagaaacag tgttctctga tgatgatttc
                                                                   4080
aatattgagc cgcttatttc tgcattccac aaagttgaga aatttgctaa ggaactgcaa
                                                                   4140
ggcaaaacag acaatcaaaa tgaccctgaa ggggaccaag aaaat
                                                                    4185
```

```
<210> 2
<211> 1404
<212> PRT
<213> Homo sapiens
```

## <400> 2

Ser Gly Ala Gln Val Lys Pro Glu Lys Lys Pro Gly Glu Glu Val Val 2.0 25 30 Gly Gly Ala Glu Ile Glu Asn Asp Val Pro Leu Val Val Arg Pro Lys 40 45 Val Arg Thr Gln Ala Gln Ile Met Pro Gly Ala Arg Pro Lys Asn Lys 55 Ser Lys Val Met Pro Gly Ala Ser Thr Lys Val Glu Thr Ser Ala Val 70 75 Gly Gly Ala Arg Pro Lys Ser Lys Ala Lys Ala Ile Pro Val Ser Arg 90 85 Phe Lys Glu Glu Ala Gln Met Trp Ala Gln Pro Arg Phe Gly Ala Glu

Tyr Trp Thr Phe Phe Val Thr Cys Thr Met Thr Gly Ala Glu Ile Glu

Arg Leu Ser Lys Thr Glu Arg Asn Ser Gln Thr Asn Ile Ile Ala Ser Pro Leu Val Ser Thr Asp Ser Val Leu Val Ala Lys Thr Lys Tyr Leu Ser Glu Asp Arg Glu Leu Val Asn Thr Asp Thr Glu Ser Phe Pro Arg Arg Lys Ala His Tyr Gln Ala Gly Phe Gln Pro Ser Phe Arg Ser Lys Glu Glu Thr Asn Met Gly Ser Trp Cys Cys Pro Arg Pro Thr Ser Lys Gln Glu Ala Ser Pro Asn Ser Asp Phe Lys Trp Val Asp Lys Ser Val Ser Ser Leu Phe Trp Ser Gly Asp Glu Val Thr Ala Lys Phe His Pro Gly Asn Arg Val Lys Asp Ser Asn Arg Ser Met His Met Ala Asn Gln Glu Ala Asn Thr Met Ser Arg Ser Gln Thr Asn Gln Glu Leu Tyr Ile Ala Ser Ser Ser Gly Ser Glu Asp Glu Ser Val Lys Thr Pro Trp Phe Trp Ala Arg Asp Lys Thr Asn Thr Trp Ser Gly Pro Arg Glu Asp Pro Asn Ser Arg Ser Arg Phe Arg Ser Lys Lys Glu Val Tyr Val Glu Ser Ser Ser Gly Ser Glu His Glu Asp His Leu Glu Ser Trp Phe Gly Ala Gly Lys Glu Gly Lys Phe Arg Ser Lys Met Arg Ala Gly Lys Glu Ala Asn Asn Arg Ala Arg His Arg Ala Lys Arg Glu Ala Cys Ile Asp Phe Met Pro Gly Ser Ile Asp Val Ile Lys Lys Glu Ser Cys Phe Trp Pro Glu Glu Asn Ala Asn Thr Phe Ser Arg Pro Met Ile Lys Lys Glu Ala Arg Ala Arg Ala Met Thr Lys Glu Glu Ala Lys Thr Lys Ala Arg Ala Arg Ala Lys Gln Glu Ala Arg Ser Glu Glu Glu Ala Leu Ile Gly Thr Trp Phe Trp Ala Thr Asp Glu Ser Ser Met Ala Asp Glu Ala Ser Ile Glu Ser Ser Leu Gln Val Glu Asp Glu Ser Ile Ile Gly Ser Trp Phe Trp Thr Glu Glu Glu Ala Ser Met Gly Thr Gly Ala Ser Ser Lys Ser Arg Pro Arg Thr Asp Gly Glu Arg Ile Gly Asp Ser Leu Phe Gly Ala Arg Glu Lys Thr Ser Met Lys Thr Gly Ala Glu Ala Thr Ser Glu Ser Ile Leu Ala Ala Asp Asp Glu Gln Val Ile Ile Gly Ser Trp Phe Trp Ala Gly Glu Glu Val Asn Gln Glu Ala Glu Glu Thr Ile Phe Gly Ser Trp Phe Trp Val Ile Asp Ala Ala Ser Val Glu Ser Gly Val Gly Val Ser Cys Glu Ser Arg Thr Arg Ser Glu Glu Glu Val Ile Gly 

```
Pro Trp Phe Trp Ser Gly Glu Gln Val Asp Ile Glu Ala Gly Ile Gly
            565
                            570
Glu Glu Ala Arg Pro Gly Ala Glu Glu Glu Thr Ile Phe Gly Ser Trp
                 585
Phe Trp Ala Glu Asn Gln Thr Tyr Met Asp Cys Arg Ala Glu Thr Ser
            600
Cys Asp Thr Met Gln Gly Ala Glu Glu Glu Pro Ile Ile Gly Ser
                  615 620
Trp Phe Trp Thr Arg Val Glu Ala Cys Val Glu Gly Asp Val Asn Ser
       630
                                635
Lys Ser Ser Leu Glu Asp Lys Glu Glu Ala Met Ile Pro Cys Phe Gly
      645 650 655
Ala Lys Glu Glu Val Ser Met Lys His Gly Thr Gly Val Arg Cys Arg
               665
        660
Phe Met Ala Gly Ala Glu Glu Thr Asn Asn Lys Ser Cys Phe Trp Ala
           680
Glu Lys Glu Pro Cys Met Tyr Pro Ala Gly Gly Gly Ser Trp Lys Ser
                   695
                                    700
Arg Pro Glu Glu Glu Glu Asp Ile Val Asn Ser Trp Phe Trp Ser Arg
       710
                                715
Lys Tyr Thr Lys Pro Glu Ala Ile Ile Gly Ser Trp Leu Trp Ala Thr
                            730
            725
Glu Glu Ser Asn Ile Asp Gly Thr Gly Glu Lys Ala Lys Leu Leu Thr
                745
Glu Glu Glu Thr Ile Ile Asn Ser Trp Phe Trp Lys Glu Asp Glu Ala
                      760
Ile Ser Glu Ala Thr Asp Arg Glu Glu Ser Arg Pro Glu Ala Glu Glu
                   775
                                   780
Gly Asp Ile Val Gly Ser Trp Phe Trp Ala Gly Glu Glu Asp Arg Leu
               790
                                795
Glu Pro Ala Ala Glu Thr Arg Glu Glu Asp Arg Leu Ala Ala Glu Lys
                            810
            805
Glu Gly Ile Val Gly Ser Trp Phe Gly Ala Arg Glu Glu Thr Ile Arg
         820 825
Arg Glu Ala Gly Ser Cys Ser Lys Ser Pro Lys Ala Glu Glu Glu
           840 845
Glu Val Ile Ile Gly Ser Trp Phe Trp Glu Glu Glu Ala Ser Pro Glu
                   855
                                   860
Ala Val Ala Gly Val Gly Phe Glu Ser Lys Pro Gly Thr Glu Glu Glu
               870
                                875
Glu Ile Thr Val Gly Ser Trp Phe Trp Pro Glu Glu Glu Ala Ser Ile
            885
                            890
Gln Ala Gly Ser Gln Ala Val Glu Glu Met Glu Ser Glu Thr Glu Glu
               905 910
         900
Glu Thr Ile Phe Gly Ser Trp Phe Trp Asp Gly Lys Glu Val Ser Glu
Glu Ala Gly Pro Cys Cys Val Ser Lys Pro Glu Asp Asp Glu Glu Met
                   935
                                   940
Ile Val Glu Ser Trp Phe Trp Ser Arg Asp Lys Ala Ile Lys Glu Thr
               950
                                955
Gly Thr Val Ala Thr Cys Glu Ser Lys Pro Glu Asn Glu Gly Ala
                             970
Ile Val Gly Ser Trp Phe Glu Ala Glu Asp Glu Val Asp Asn Arg Thr
                         985
         980
Asp Asn Gly Ser Asn Cys Gly Ser Arg Thr Leu Ala Asp Glu Asp Glu
      995 1000
Ala Ile Val Gly Ser Trp Phe Trp Ala Gly Asp Glu Ala His Phe
```

	1010					1015					1020			
Glu		Asn	Pro	Ser	Pro		Phe	Arg	Ala	Ile	Cys	Arg	Ser	Thr
	1025					1030		_			1035	_		
Cys	Ser	Val	Glu	Gln	Glu	Pro	Asp	Pro	Ser	Arg	Arg	Pro	Gln	Ser
	1040					1045					1050			
Trp	Glu	Glu	Val	Thr	Val	Gln	Phe	Lys	Pro	Gly	Pro	Trp	Gly	Arg
	1055					1060					1065			
Val	_	Phe	Pro	Ser	Ile	Ser	Pro	Phe	Arg	Phe	Pro	rys	Glu	Ala
	1070					1075					1080			
Ala		Leu	Phe	Суѕ	Glu		Phe	Gly	Gly	Lys	Pro	Arg	Asn	Met
77-7	1085	G	D	G1	C1	1090	7	C1	C1	G	1095	T	C1	D
vai	Leu 1100	ser	PIO	GIU	GIY	1105	Asp	GIII	GIU	ser	Leu 1110	ьеu	Gln	FIO
Asp		Pro	Ser	Pro	Glu		Pro	Phe	Gln	Tvr	Asp	Pro	Ser	Tvr
1	1115	_ =		_ =		1120	_ =				1125	_ =		
Arg	Ser	Val	Gln	Glu	Ile	Arg	Glu	His	Leu	Arg	Ala	Lys	Glu	Ser
	1130					1135					1140			
Thr	Glu	Pro	Glu	Ser	Ser	Ser	Cys	Asn	CAa	Ile	Gln	CA2	Glu	Leu
	1145					1150					1155			
Lys		Gly	Ser	Glu	Glu		Glu	Glu	Leu	Leu	Leu	Leu	Met	Glu
-	1160		-	_	D.I	1165		a i	- 1	<b>a</b>	1170	- 1		
гуз	11e	Arg	Asp	Pro	Pne	11e	HIS	GIU	TTe	ser	Lys 1185	IIe	Ala	мес
Glv		Ara	Ser	Ala	Ser		Phe	Thr	Ara	Asp	Phe	Tle	Ara	Asp
1	1190	9				1195			9	1-	1200		9	
Ser	Gly	Val	Val	Ser	Leu	Ile	Glu	Thr	Leu	Leu	Asn	Tyr	Pro	Ser
	1205					1210					1215			
Ser	Arg	Val	Arg	Thr	Ser		Leu	Glu	Asn	Met	Ile	Arg	Met	Ala
_	1220	_	_	_	_	1225	_ •	_ •			1230		_	_
Pro	1235	Tyr	Pro	Asn	Leu	Asn 1240	lle	lle	GIn	Thr	Tyr 1245	lle	Cys	Lys
Val		Glu	Glu	Thr	T.e.11		Tur	Ser	Val	Asn	Ser	Pro	Glu	Gln
	1250	-14	-14			1255	- <u>v</u> -				1260			
Leu		Gly	Ile	Arg	Met		Arg	His	Leu	Thr	Thr	Thr	Thr	Asp
	1265					1270					1275			
Tyr	His	Thr	Leu	Val	Ala	Asn	Tyr	Met	Ser	Gly	Phe	Leu	Ser	Leu
	1280					1285					1290			
Leu		Thr	Gly	Asn	Ala	-	Thr	Arg	Phe	His	Val	Leu	Lys	Met
T .	1295	7	T.	<b>G</b> .	a i	1300	<b>T</b> .	D.I	N/ - !-	m1	1305	<i>a</i> 3	<b>T</b> .	т.
ьеи	Leu 1310	ASN	ьeu	ser	GIU	Asn 1315	ьeu	rne	met	ınr	Lys 1320	GIU	ьeu	Leu
Ser		Glu	Ala	Val	Ser		Phe	Ile	Glv	Leu	Phe	Asn	Ara	Glu
201	1325	o_u			~~_	1330			<u> y</u>		1335		9	014
Glu		Asn	Asp	Asn	Ile		Ile	Val	Leu	Ala	Ile	Phe	Glu	Asn
	1340					1345					1350			
Ile	Gly	Asn	Asn	Ile	Lys	Lys	Glu	Thr	Val	Phe	Ser	Asp	Asp	Asp
	1355					1360					1365			
Phe		Ile	Glu	Pro	Leu		Ser	Ala	Phe	His	Lys	Val	Glu	Lys
Db-	1370	T	C1	T ~ · ·	C1	1375	T				1380			
rne	Ala	тЛа	GIU	ьeu	GTII	ату	тЛа							